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SEQUENCE LISTING

<110> THE JOHNS HOPKINS UNIVERSITY SCHOOL OF MEDICINE
PEDERSEN, Peter L.
MATHUPALA, Saroj P.

<120>   ARREST OF PROLIFEATION OF HIGHLY GLYCOLYTIC TUMORS

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atatttgaat gtatttagaa aaataaacaa ataggggttc cgcgcacatt tccccgaaaa    6360

gtgccacctg acgtctaaga aaccattatt atcatgacat taacctataa aaataggcgt    6420

atcacgaggc cctttcgtct tcaa                                           6444
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<213>  Artificial sequence

<220>
<223>  Forward primer

<400>  4
ggaattcatg atcgcctcgc atatgatc                                        28


<210>  5
<211>  29
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<212>  DNA
<213>  Artificial sequence

<220>
<223>  Revers primer

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gctctagact atctctgccc agcctcccg                                        29


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<213>  Artificial sequence

<220>
<223>  Forward primer

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cggatccgcc cctctccctc cc                                               22


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<211>  24
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<220>
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gatagcggtc cgccacaccc agcc                                             24


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<212>  DNA
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<220>
<223>  Forward primer

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<212>  PRT
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<213> Rattus norvegicus

<400> 10

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Met Ile Ala Ser His Met Ile Ala Cys Leu Phe Thr Glu Leu Asn Gln
1               5                   10                  15

Asn Gln Val Gln Lys Val Asp Gln Phe Leu Tyr His Met Arg Leu Ser
            20                  25                  30

Asp Glu Thr Leu Leu Glu Ile Ser Arg Arg Phe Arg Lys Glu Met Glu
        35                  40                  45

Lys Gly Leu Gly Ala Thr Thr His Pro Thr Ala Ala Val Lys Met Leu
    50                  55                  60

Pro Thr Phe Val Arg Ser Thr Pro Asp Gly Thr Glu His Gly Glu Phe
65                  70                  75                  80

Leu Ala Leu Asp Leu Gly Gly Thr Asn Phe Arg Val Leu Arg Val Arg
            85                  90                  95

Val Thr Asp Asn Gly Leu Gln Arg Val Glu Met Glu Asn Gln Ile Tyr
            100                 105                 110

Ala Ile Leu Glu Asp Ile Met Arg Gly Ser Gly Thr Gln Leu Phe Asp
        115                 120                 125

His Ile Ala Glu Cys Leu Ala Asn Phe Met Asp Lys Leu Gln Ile Lys
    130                 135                 140

Glu Lys Lys Leu Pro Leu Gly Phe Thr Phe Ser Phe Pro Cys His Gln
145                 150                 155                 160

Thr Lys Leu Asp Glu Ser Phe Leu Val Ser Trp Thr Lys Gly Phe Lys
            165                 170                 175

Ser Ser Gly Val Glu Gly Arg Asp Val Val Asp Leu Ile Arg Lys Val
            180                 185                 190

Ile Gln Arg Arg Gly Asp Phe Asp Ile Asp Ile Val Ala Val Val Asn
        195                 200                 205

Asp Thr Val Gly Thr Met Met Thr Cys Gly Tyr Asp Asp Gln Asn Cys
    210                 215                 220

Glu Ile Gly Leu Ile Val Gly Thr Gly Ser Asn Ala Cys Tyr Met Glu
225                 230                 235                 240

Glu Met Arg His Ile Asp Met Val Glu Gly Asp Glu Gly Arg Met Cys
            245                 250                 255

Ile Asn Met Glu Trp Gly Ala Phe Gly Asp Asp Gly Thr Leu Asn Asp
            260                 265                 270

Ile Arg Thr Glu Phe Asp Arg Glu Ile Asp Met Gly Ser Leu Asn Pro
        275                 280                 285

Gly Lys Gln Leu Phe Glu Lys Met Ile Ser Gly Met Tyr Met Gly Glu
    290                 295                 300
```

Leu Val Arg Leu Ile Leu Val Lys Met Ala Lys Ala Glu Leu Leu Phe
305   310   315   320

Gln Gly Lys Leu Ser Pro Glu Leu Leu Thr Thr Gly Ser Phe Glu Thr
                325   330   335

Lys Asp Val Ser Asp Ile Glu Glu Asp Lys Asp Gly Ile Glu Lys Ala
            340   345   350

Tyr Gln Ile Leu Met Arg Leu Gly Leu Asn Pro Leu Gln Glu Asp Cys
        355   360   365

Val Ala Thr His Arg Ile Cys Gln Ile Val Ser Thr Arg Ser Ala Ser
    370   375   380

Leu Cys Ala Ala Thr Leu Ala Ala Val Leu Trp Arg Ile Lys Glu Asn
385   390   395   400

Lys Gly Glu Glu Arg Leu Arg Ser Thr Ile Gly Val Asp Gly Ser Val
            405   410   415

Tyr Lys Lys His Pro His Phe Ala Lys Arg Leu His Lys Ala Val Arg
            420   425   430

Arg Leu Val Pro Asp Cys Asp Val Arg Phe Leu Arg Ser Glu Asp Gly
        435   440   445

Ser Gly Lys Gly Ala Ala Met Val Thr Ala Val Ala Tyr Arg Leu Ala
    450   455   460

Asp Gln His Arg Ala Arg Gln Lys Thr Leu Glu Ser Leu Lys Leu Ser
465   470   475   480

His Glu Gln Leu Leu Glu Val Lys Arg Arg Met Lys Val Glu Met Glu
            485   490   495

Gln Gly Leu Ser Lys Glu Thr His Ala Val Ala Pro Val Lys Met Leu
            500   505   510

Pro Thr Tyr Val Cys Ala Thr Pro Asp Gly Thr Glu Lys Gly Asp Phe
        515   520   525

Leu Ala Leu Asp Leu Gly Gly Thr Asn Phe Arg Val Leu Leu Val Arg
    530   535   540

Val Arg Asn Gly Lys Arg Arg Gly Val Glu Met His Asn Lys Ile Tyr
545   550   555   560

Ser Ile Pro Gln Glu Val Met His Gly Thr Gly Glu Glu Leu Phe Asp
            565   570   575

His Ile Val Gln Cys Ile Ala Asp Phe Leu Glu Tyr Met Gly Met Lys
            580   585   590

Gly Val Ser Leu Pro Leu Gly Phe Thr Phe Ser Phe Pro Cys Gln Gln
            595   600   605

Asn Ser Leu Asp Gln Ser Ile Leu Leu Lys Trp Thr Lys Gly Phe Lys
610   615   620

Ala Ser Gly Cys Glu Gly Glu Asp Val Val Thr Leu Leu Lys Glu Ala

```
              625                      ·630                    635                      640

              Ile His Arg Arg Glu Glu Phe Asp Leu Asp Val Val Ala Val Val Asn
                          645             650                 655

              Asp Thr Val Gly Thr Met Met Thr Cys Gly Tyr Glu Asp Pro His Cys
                          660             665                 670

              Glu Val Gly Leu Ile Val Gly Thr Gly Ser Asn Ala Cys Tyr Met Glu
                          675             680                 685

              Glu Met Arg Asn Val Glu Leu Val Asp Gly Glu Glu Gly Arg Met Cys
                  690             695                 700

              Val Asn Met Glu Trp Gly Ala Phe Gly Asp Asn Gly Cys Leu Asp Asp
              705             710                 715                     720

              Leu Arg Thr Val Phe Asp Val Ala Val Asp Glu Leu Ser Leu Asn Pro
                          725             730                 735

              Gly Lys Gln Arg Phe Glu Lys Met Ile Ser Gly Met Tyr Leu Gly Glu
                          740             745                 750

              Ile Val Arg Asn Ile Leu Ile Asp Phe Thr Lys Arg Gly Leu Leu Phe
                          755             760                 765

              Arg Gly Arg Ile Ser Glu Arg Leu Lys Thr Arg Gly Ile Ser Glu Thr
                  770             775                 780

              Lys Phe Leu Ser Gln Ile Glu Ser Asp Cys Leu Ala Leu Leu Gln Val
              785             790                 795                     800

              Arg Ala Ile Leu Arg His Leu Gly Leu Glu Ser Thr Cys Asp Asp Ser
                          805             810                 815

              Ile Ile Val Lys Glu Val Cys Thr Val Val Ala Arg Arg Ala Ala Gln
                          820             825                 830

              Leu Cys Gly Ala Gly Met Ala Ala Val Val Asp Lys Ile Arg Glu Asn
                          835             840                 845

              Arg Gly Leu Asp Asn Pro Lys Val Thr Val Gly Val Asp Gly Thr Leu
                  850             855                 860

              Tyr Lys Leu His Pro His Phe Ala Lys Val Met His Glu Thr Val Arg
              865             870                 875                     880

              Asp Leu Ala Pro Lys Cys Asp Val Ser Phe Leu Glu Ser Glu Asp Gly
                          885             890                 895

              Ser Gly Lys Gly Ala Ala Leu Ile Thr Ala Val Ala Cys Arg Ile Arg
                          900             905                 910

              Glu Ala Gly Gln Arg
                          915
```